Synthesis Request Sheet () Apple G 497810 WILLITY USE ONLY: Sequence # 3242 5 Acceled G 497852 Column Date Synthesized _ | | Sequence Length: 36
Sequestor: Gersten Synthesis Scale: 0.2 umole
F.i. Name: LOWE Trityl Group: __On _X_ Trityl Group: ___On _X_ Off Bi:ling Dept.: HHMI Thio-Chemistry: ___Yes _X_ No Shone Number: 7-4756 Account Number: 921099 SEQUENCE. 5'-) 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *) 1908 DIT 194 GEA AAC AGG AAG 9AC AGG AGG ETC TGG User Comments: FucI-VII genomic por bp 9-34 5' por exon 1 Comments to Core Facility: - 194 Synthesis Setup Listing - (Version 1.02) Islamn 1 Pun ID : Ivole : Ovc01 user End Proc: End CE DMT = Off) Sequence: 32428 Total bases = 36 A= 10. G= 13. C= 10. T= 3, 5= 0, 6= 0. 7= 0. 8= 0 (mixed bases= 0) MW: 11142.2

Exhibit C

5'> GCG CCT CGA GCA AAC AGG AAG GAC AGG AGG CTC TGG <3'

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

1088 B

Column

Appled 425287 电多多分类 医多种性性 医皮肤 医三二二

Date Synthesized __

Today's Date:

Requestor: Gersten F.I. Name: LOWE

Sequence Length: 20

Synthesis Scale: 0.2 umole

Billing Dept.: HHMI

Trityl Group: ___On _X_ Off
Thio-Chemistry: ___Yes _X_ No

Phone Number: 7-4754 Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

TTT GTT TCC GCC CCG GCA TC

User Comments:

104 phage from primer 1042 towards sal site

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 3

1. The

#2.

Run ID :

Cycle : Cyc01 user

End Proc: End CE

(DMT = Off)

Sequence: 1088B

Total bases = 20

A= 1, G= 4, C= 8, T= 7, S= 0, G= 0, T= 0, T= 0

(mixed bases= 0)

MW: 6418.0

✓5'> TIT GIT TCC GCC CCG GCA TC <3'

A200 - .440

42 ml /958

189928 Synthesis Request Sheet ------Sequence # FACILITY USE ONLY: Applied C 483082 Column ------Date Synthesized ____ Sequence Length: 20 Today's Date: Requestor: Gersten
P.I. Name: LOWE
Billing Dept.: HHMI
Phone Number: 7-4756 Synthesis Scale: 0.2 umole Trityl Group: __On _X_ Off
Thio-Chemistry: __Yes _X_ No Account Number: 030131 SEQUENCE, 5'-> 3'. IN TRIPLETS (* PLEASE START WITH A TRIPLET *) GCT AGA TAG TTT CTG ATG GC OD240 1,00= ,14/ User Comments: Fuct7-1 Comments to Core Facility: - 394 Synthesis Setup Listing -(Version 2.00) Column 2 Run ID : Cycle : Cyc01 user (DMT = Off)End Proc: End CE

Sequence: 9392A

Total bases = 20 A= 4,6= 6,C= 3,T= 7,5= 0,6= 0,7= 0,8= 0 (mixed bases= 0)

MW: 6158.0

5'> GCT AGA TAG TTT CTG ATG GC <3'

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

Column

Acceled C 436432

Date Synthesized _____

Today's Date:

Sequence Length: 20

Requestor:

Gersten LOWE

Synthesis Scale: 0.2 umole

Trityl Group: __On _X_ Off
Thio-Chemistry: __Yes _X_ No

P.I. Name:

Billing Dept.: HHMI Phone Number:

7-4754

Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

GTA TGG GTG CCA TCA GAA AC

User Comments:

104 sal-eco from 986 towards 5' UT

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 4

Run ID :

Cycle : Cyc01 user

End Proc: End CE (OMT = Off)

Sequence: 1042B

Total bases = 20

A= 6, G= 6, C= 4, T= 4, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 6654.0

5'> GTA TGG GTG CCA TCA GAA AC <3'

Azco = .217

46/954

STOURD TORE RESEMPTION CORE PAULILITES DNA SYNTHESIS CORE 2562 MSRB II 764-1461

Synthesis Request Sheet

CILITY USE ONLY:

Sequence #

Column

Date Synthesized

Today's Date:

Gersten

Sequence Length: 20

Requestor: P.I. Name: LOWE

Synthesis Scale: 0.2 umole

Trityl Group: ___On _X_ Off
Thio-Chemistry: ___Yes _X_ No

Billing Dept.: HHMI Phone Number: 7-4756 Account Number: 030131

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

CAG GGC ACT TCT GAG GTG CC

User Comments:

Fuc-TVII bp424 sequencing primer towards 3' end

Comments to Core Facility:

1/100 - .405

- 394 Synthesis Setup Listing - (Version 1.01)

Column 3

Run ID :

Cycle : Cyc01 user

End froc: End CE

(DMT = Off)

Sequence: 8660A

Total bases = 20

A= 3, G= 7, C= 6, T= 4, 5= 0, 6= 0, 7= 0, 8= 0 (mixed bases= 0)

MW: 6573.0

5'> CAG GGC ACT TCT GAG GTG CC <3'

© 495751 Synthesis Request Sheet FACILITY USE ONLY: Sequence # Appled G 472563 Column D.6. Date Synthesized ___ Today's Date: Sequence Length: 20 Requestor: Gersten Synthesis Scale: 0.2 umole P.I. Name: LOWE Trityl Group: __On _X_ Off
Thio-Chemistry: __Yes _X_ No Billing Dept.: HHMI Phone Number: 7-4756 Account Number: 030131 SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *) AAC AGC AGC CTT GTC ACG TG 00 260 1100 = .156 User Comments: Fuct7-2 Comments to Core Facility: - 394 Synthesis Setup Listing - (Version 2.00) Column 3 Run ID : Cycle : Cyc01 user End Proc: End CE (DMT = Off) Sequence: 9393A Total bases = 20 A= 5, G= 5, C= 6, T= 4, 5= 0, 6= 0, 7= 0, 8= 0 (mixed bases≃ 0) MW: 6078.0

5'> AAC AGC AGC CTT GTC ACG TG <3'

Synthesis Request Sheet

| FACILITY USE ONLY: | Sequence # | |
|--|--|-------------------------|
| | Column | © 121275 |
| | Date Synthesized | <u></u> ,(B) |
| Today's Date: Requestor: Gersten P.I. Name: LOWE Billing Dept.: HHMI Phone Number: 7-4778 Account Number: 921099 | Sequence Length: Synthesis Scale: Trityl Group: Thio-Chemistry: | O.2 umole On _X_ Off |
| SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *) | | |
| CAG CCA TAG TCT CAC GTG AC | | |
| User Comments: 104 sal-eco from 887 primer towards sal site (5' UT) | | |
| Comments to Core Facilit394 Synthe | esis Setup Listing - | (Version 1.01) |
| Column 3 | | |
| Run ID : Cycle : Cyc01 user End Proc: End CE (DMT = Off) Sequence: 986B | , | |
| Total bases = 20 A= 5, G= 4, C= 7, T= (mixed bases= 0) | 4, 5= 0, 6= 0, 7= | 0, 8= 0 |

5'> CAG CCA TAG TCT CAC GTG AC (3'

MW: 6536.0

Synthesis Request Sheet

FACILITY USE ONLY: Sequence # 672 B Column Today's Date:

Requestor:

Bersten

F.I. Name:

LOWE

Billing Dept.: Trityl Group: __On _X_ Off Thio-Chemistry: __Yes Y_ No Phone Number: 7-4778 Account Number: 921079 SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *) CGA AGT GTA GGA AGT GAT CC User Comments: 104 from 15/52 large (Xho-Kph) towards 3' end Comments to Core Facility: - 394 Synthesis Setup Listing - (Version 1.01) grand and the state of the stat Column 2 Run ID : Cycle : Cyc01 user End Proc: End CE (DMT = Off) Sequence: 6728 Total bases = 20 A= 6, G= 7, C= 3, T= 4, 5= 0, 6= 0, 7= 0, 8= 0(mixed bases= 0) MW: 6693.0

-260 = ,304

√ 5'> CGA AGT GTA GGA AGT GAT CC <3'

33 N 1967

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

887 B

Column

Accepted C 419751

Date Synthesized ___ _____

Today's Date:

Sequence Length: 20

Requestor: Gersten
P.I. Name: LOWE

Synthesis Scale: 0.2 umole

Trityl Group: ___On _X_ Off
Thio-Chemistry: ___Yes _X_ No

Billing Dept.: HHMI Phone Number: 7-4778 Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

CAC TCT GCT GAG CCA GAC AC

User Comments:

104 sequencing from sac-T7 towards upstream exons

Comments to Core Facility:

COLUMN I SET-UP VERSION 2.01

USER_NAME:

CYCLES USED:

SSCEAF36- I

ENDING METHOD:

Trityl OFF, Auto

ENDING PROCEDURE: DEPCRE03

SEQUENCE NAME:

887B

SEQUENCE LENGTH: 20

DATE: TIME:

COMMENT:

V5'- CAC TOT GOT GAG CCA GAC AC -3'

Az60-2 -289

Report for Synthetic Oligonucleotide UPTH FREE SINGS SI Date:

Investigator: LOWE

Individual User: Gersten

System Id: 8993

Product Information

Length: 20

Oligonucleotide Sequence: AGG AAG CTT AGC TAA AAG CT

Comments:

104 phage from t7 primer towards 3' end of gene (5' ut towards gene)

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 2

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8993 Thank you for using the HHMI Biopolymer Facility.

Use 104 Sac PTZF ONA

A260 = . 264

38 Wl /962

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

Column

Acoled C 410554 **以及以及其他是是是是是其他的**的。

Date Synthesized _____

Sequence Length: 20 Synthesis Scale: 0.2 umole

Today's Date:
Requestor: Gersten
P.I. Name: LOWE
Billing Dept.: HHMI
Phone Number: 7-4778

Trity1 Group: __On _X Off Thio-Chemistry: __Yes _3 No

Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET 4)

GCA AAG CTA TAG CTT GTA GC

User Comments:

104 from15/52(EcoRI) towards 5'end

Comments to Core Facility:

- 394 Synthesis Setup Listing -

(Version 1.01)

* 4: . x

Column 1

Run ID :

Cycle : Cyc01 user

End Proc: End CE (DMT = Off)

Sequence: 671B

Total bases = 20

A= 6, G= 5, C= 4, T= 5, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 6629.0

5'> GCA AAG CTA TAG CTT GTA GC <3'

Aug = .319

31 wl / 969

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

3511 B

Appled G 404333

Column

Date Synthesized

Date Submitted:

Kelly

Sequence Length: 20

Requestor: Phone Number: 74756

Synthesis Scale: 0.2 umole

Account Number: 921099

Trityl Group:

No

P.I. Name: LOWE

HPLC Purify:

Billing Dept.: HHMI

Thio-Chemistry: No

Center Membership: None

Center/Project Related Research: No

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

AGC CTG GAC CTG AGG CTG GG

User Comments:

FT7-1

Comments to Core Facility:

- 394 Synthesis Setup Listing -

(Version 2.00)

Column 2

Run ID :

Cvcle : Cvc01 user

End Proc: End CE

(DMT = Dff)

Sequence: 35118

Total bases = 20

A= 3, G= 9, C= 5, T= 3, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 6191.0

5'> AGC CTG GAC CTG AGG CTG GG <3'

0D20 1/00 .102 40/ml

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

946 B

Column

Appled G 439483

Date Synthesized

D.G.

Sequence Length: 20

Today's Date:
Requestor: Gersten P.I. Name:

LOWE

Synthesis Scale: 0.2 umole

Billing Dept.: HHMI

Trityl Group: ___On _X_ Off
Thio-Chemistry: ___Yes _X_ No

Phone Number: 7-4778 Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

TCC TTC CCT TTC CAG ACT GG

User Comments:

104 sac sequencing from 910 towards cat exon

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 4

Run ID :

Cycle : Cyc01 user

End Proc: End CE (DMT = Off)

Sequence: 946B

Total bases = 20

A= 2, G= 3, C= 8, T= 7, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 6419.0

√5"> TCC TTC CCT TTC CAG ACT GG (3"

A260 = ,204

NEW TOTAL TANK THE PROPERTY OF THE PARTY OF DNA SYNTHESIS CORE 2562 MSRB II 764-1461

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

715 B

Column

Acceptant C 406550

Date Synthesized __

Todav's Date:

Sequence Length: 20

Requestor: Gersten P.I. Name: LOWE

Synthesis Scale: 0.2 umole

Trityl Group: __On _X Off
Thio-Chemistry: __Yes _X No

Billing Dept.: HHMI Fhone Number: 7-4778

Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

GGG AAG GAG TOT GTG TOT CC

User Comments:

104 per of t cell large frag from sp6 seq

Comments to Core Facility:

COLUMN Z SET-UP VERSION Z.01

USER NAME:

CYCLES USED:

SSCEAF3G- I

ENDING METHOD: Trityl OFF, Auto

ENDING PROCEDURE: DEPCRE03

SEQUENCE NAME:

715B

SEQUENCE LENGTH: 20

DATE: TIME:

COMMENT:

√5'- GGG AAG GAB TCT GTG TGT CC -3'

A260 = ,304

33 ul /967

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence # 2931 B

Column

do Applied G 479590

Date Synthesized _____ Clin Belle

Today's Date:

Requestor: Gersten
P.I. Name: LOWE

Billing Dept.: HHMI Phone Number: 7-4756

Account Number: 921099

Sequence Length: 20

Synthesis Scale: 0.2 umole

Trityl Group: ___On _X_ Off
Thio-Chemistry: ___Yes _X_ No

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

ACC TTG GGC TGA ACC TAC AG

User Comments:

sequence primer from 946 towards exon 1 S.D. (FT-VII)

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 3

Run ID :

Cycle : Cyc01 user

End Proc: End CE

(DMT = Off)

Sequence: 2931B

Total bases = 20

A= 5, G= 5, C= 6, T= 4, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= ⊙)

MW: 6575.0

5'> ACC TTG GGC TGA ACC TAC AG <3'

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

2932 B

Column

Appled C 460602

Date Synthesized __

Today's Date:

Sequence Length: 20

Requestor: Gersten
P.I. Name: LOWE

Synthesis Scale: 0.2 umole

Billing Dept.: HHMI

Trityl Group: __On _X_ Off
Thio-Chemistry: __Yes _X_ No

Phone Number: 7-4756 Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

User Comments:

sequence primer proximal to exon 1 S.D towards 5' UT (FT-VII)

Comments to Core Facility:

CCT TGA ACT GTA GGT TCA GC

- 394 Synthesis Setup Listing - (Version 1.01)

Column 2

Run ID :

Cycle : Cyc01 user

End Proc: End CE (DMT = Off)

Sequence: 2932B

Total bases = 20

A= 4, G= 5, C= 5, T= 6, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= ())

MW: 6563.0

5'> CCT TGA ACT GTA GGT TCA GC <3'

Synthesis Request Sheet

CILITY USE ONLY:

Sequence #

Column

C 184064

Date Synthesized ____

Today's Date:

Gersten

Sequence Length: 20

Requestor:

LOWE

Synthesis Scale: 0.2 umole

P.I. Name: Billing Dept.: HHMI

Trityl Group:

Phone Number: 7-4756

Trityl Group: ___On _X_ Off
Thio-Chemistry: ___Yes _X_ No

Account Number: 030131

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

TTT TCT AGA GGT GGC AGA AC

User Comments:

fuc-tvii sequencing primer towards 3' end (bp 1333)

1/100 =

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 3

Run ID :

Cycle : Cyc01 user

End Proc: End CE

(DMT = Off)

Sequence: 8661A

Total bases = 20

A= 5, G= 6, C= 3, T= 6, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 6642.0

5'> TTT TCT AGA GGT GGC AGA AC <3'

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

911 B

Column

Appled C 426226 ----

Date Synthesized __

Today's Date:

Sequence Length: 20

Synthesis Scale: 0.2 umole

Requestor: Gersten
P.I. Name: LOWE

Billing Dept.: HHMI Phone Number: 7-4778

Trityl Group: __On _X Off
Thio-Chemistry: __Yes _X No

Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

GGA CCT GTG CTC CCA GAT CC

User Comments:

104 sequencing from sac 9076 primer towards 5' exon

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 2

Run ID :

Cycle : Cyc01 user

End Proc: End CE

Sequence: 911B

Total bases = 20

A= 3, G= 5, C= 8, T= 4, 5= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 6495.0

√5'> GGA CCT GTG CTC CCA GAT CC <3'

Aug 2.215

47 1 /953

Report for Synthetic Oligonucleotide 100 this total him was been seen to 100 and the cold and cold and the cold that the cold and the

Date:

Investigator: LOWE

Individual User: Gersten

System Id: 9077

Product Information

Length: 20

Oligonucleotide Sequence: AAG GGA TCA CTT CTG CTC AG

Comments:

104phage from 8952 towards 5' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 4

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 9077 Thank you for using the HHMI Biopolymer Facility.

Azao = .219

46 W / 954

Report for Synthetic Oligonucleotide And also such that man had man and that may and and that may and and man and man and the plant and the first man in the plant in a dark

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

9076

Product Information

Length: 20

Oligonucleotide Sequence: TGC TT(CAGT) CCT TCA GGA AAA GG

Comments:

104phage from 8952 towards 5' ut

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 9076 Thank you for using the HHMI Biopolymer Facility.

H20 = ,217

461954

Synthesis Request Sheet

FACILITY USE ONLY:

3445 B

Applied G 404852

Column

Date Synthesized

Date Submitted:

Kelly

Sequence Length: 20 Synthesis Scale: 0.2 umole

Requestor: Kelly Phone Number: 74756

Account Number: 921099 P.I. Name: LOWE

Trityl Group: HPLC Purify:

No

Billing Dept.: HHMI

Thio-Chemistry: No

Center Membership: None

Center/Project Related Research: No

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

CTA GCT GGT CAT TTC TAG GG

User Comments:

ft7-1

Comments to Core Facility:

DNA SEQUENCE 1

NUMBER OF BASES: 20

BASES USED: A=3 C=4 6=6 T=7 X=0

DALTONS:

5115

TIME:

DATE:

E' > OTA GOT SGT CAT TTC TAG GG < 3'

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8953

Product Information

Length: 20

Oligonucleotide Sequence: TCA ATT CCC TCT TTG AGC AG

Comments:

104 phage from 8903 primer towards 3' end

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8953 Thank you for using the HHMI Biopolymer Facility.

A260 = , 290

38 ml / 965 ml

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8952

Product Information

Length: 20

Oligonucleotide Sequence: ATC AAC CAC TAT CCA ATC CT

Comments:

104 phage from primer 8903 towards 5' end

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8952 Thank you for using the HHMI Biopolymer Facility.

Ango = . 290

75al (965al

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8904

Product Information

Length: 20

Oligonucleotide Sequence: TGA CAA TTC CAG AAG GCT CC

Comments:

104 phage from primer 8874 towards 3' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8904 Thank you for using the HHMI Biopolymer Facility.

A260 = ,272

37 N /963

Date:

Investigator: LOWE

Individual User: Gersten

System Id: 8903

Product Information

Length: 20

Oligonucleotide Sequence: GGC CAG GCA CTC ACC AAT AC

Comments:

104 phage from primer 8874 towards 5' end of gene

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8903 Thank you for using the HHMI Biopolymer Facility.

A 260 = .292

35 wl /965

Date:

Investigator: LOWE

System Id: 8902

Individual User: Gersten

Product Information

Length: 20

Oligonucleotide Sequence: TTA TTC TGC TTC AGG GTA CC

Comments:

104 phage from primer 8874 towards 3' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8902 Thank you for using the HHMI Biopolymer Facility.

A260 = . 297

342 /966

Date:

Investigator: LOWE

Individual User: Gersten

System Id: 8875

Product Information

Length: 20

Oligonucleotide Sequence: ATC TGC ACT GGC CTT TCA CC

Comments:

104 phage from primer 8850 towards 3' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received:
Date Synthesised:
Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8875
Thank you for using the HHMI Biopolymer Facility.

A260 = .300

33 al (967 ml

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

524 B

Column

Appled C 416455

Oate Synthesized _

Todav's Date:

Sequence Length: 41

Requestor: Gersten

Synthesis Scale: 0.2 umole

P.I. Name:

LOWE

Billing Dept.: HHMI

Trityl Group: __On _X_ Off Thio-Chemistry: __Yes [%] No

Phone Number: 7-4778

Account Number: 921099

SEQUENCE, 5'-> 3'. IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

GCG CBG ATC CPA CCA TCC TTA TCT GBC ACT GGC CTT TCA CC

User Comments:

phage 104 FCR primer (FOR AB) from nt 282

Comments to Core Facility:

- 394 Synthesis Setup Listing - (Version 1.01)

Column 4

Run ID :

Cycle : Cyc01 user

End Proc: End CE (DMT = Off)

Sequence: 624B

Total bases = 41

A= 6, G= 8, C= 17, T= 10, S= 0, 6= 0, 7= 0, 8= 0

(mixed bases= 0)

MW: 13246.2

Buren Ill

✓ 5'> GCG QGG ATC QCA CCA TCC TTA TCT GGC ACT GGC CTT TCA

CC <3'

Az60 = .505 200 25W /15W

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8874

Product Information

Length: 20

Oligonucleotide Sequence: TGG TGA GTG TGG ACT GAG GC

Comments:

104 phage from primer 8850 towards 5' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised: Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8874 Thank you for using the HHMI Biopolymer Facility.

A240 = ,300

33 W/967

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8867

Froduct Information

Length: 20

Oligonuclectide Sequence: GCT AGC AGG CTC CGG TTA GC

Comments:

104phage from 8770primer (b sequence) towards 5' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8867
Thank you for using the HHMI Biopolymer Facility.

Az60 = .259

38 ll /962

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8851

Product Information

Length: 20

Oligonucleotide Sequence: CCT TGG GTC TGG GCC TCC AT

Comments:

104 phage from 8770 primer towards 3' end of gene

.

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 1

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8851 Thank you for using the HHMI Biopolymer Facility.

A260 - . 281

36 ul /9

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8850

Product Information

Length: 20

Oligonuclectide Sequence: AAG CGA TAG AGA CCA TGG GT

Comments:

104 phage from 8770 primer towards 5' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Received:
Date Synthesised:

Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8850 Thank you for using the HHMI Biopolymer Facility.

A260 = , 264

39 ul / 962

Date:

Investigator: LOWE

System Id: 8771

Individual User: Gersten

Product Information

the side that the property of the contract of the side of the side

Length: 20

Oligonucleotide Sequence: GGC CCA CAT CCC CAC TAC CG

Comments:

104 phage from 8715 sequence towards 3' end (towards rp2 primer seque

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8771 Thank you for using the HHMI Biopolymer Facility.

A260 = . 255

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8852

Product Information

MATERIAL BOOK SHOW HAVE A THE WARE A METERIAL BOOK SHOW SHOW THE BOOK SHOW THE

Length: 20

Oligonucleotide Sequence: CAC GCT GCT GCC GCT CCT GG

Comments:

104 phage from 8715 primer towards 5' end of gene (replace primer 8770

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 1

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8852
Thank you for using the HHMI Biopolymer Facility.

A260 = .265

38 wl /962

Report for Synthetic Oligonucleotide the less two was are 100 and and 110 and 110 and 110 and 100 a

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8714

Product Information

and one i was year the was the control of the contr

Length: 20

Oligonucleotide Sequence: GCA TCG GGA CTA CAT CAC TG

Comments:

104pst from rp2 sequence towards the 3' end

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8714 Thank you for using the HHMI Biopolymer Facility.

A260 = .266

38 w olyo 962 w H20

Report for Synthetic Oligonucleotide Miles wide and acted processing the control form the cont

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8715

Product Information

Length: 20

Oligonucleotide Sequence: AGC CCC AGG CAT TGC GCC AG

Comments:

104pst from rp2 primer towards the 5' end

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8715 Thank you for using the HHMI Biopolymer Facility.

ALGO = .297

33 ul oligio 967 ul H20

Date:

make rang pligt at a case and a play high total make and play high bear make the case of t

Investigator: LOWE

Individual User: Gersten

System Id:

8848

Product Information

Length: 20

Oligonucleotide Sequence: AAC TEE CTE TCT TCC TCE TC

Comments:

104phage from primer 8714 towards 3' end

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 1

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8848 Thank you for using the HHMI Biopolymer Facility.

Report for Synthetic Oligonucleotide The N h was the tip tip and the set and th

Date:

Investigator: LOWE

Individual User: Gersten

System Id: 8849

Product Information

traft effet atom spiret mooth erest until and a spire atom cross for an house mont force arter dates some com-

Length: 20

Oligonucleotide Sequence: CAC GAT AAC GAC TOT CAT TO

Comments:

104 phage from primer 8714 towards 5' end

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 1

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8849 Thank you for using the HHMI Biopolymer Facility.

Az60 = .283

35 W / 965

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8888

Product Information

Length: 20

Oligonucleotide Sequence: CTG GAG GGA AGC GCT TCT GC

Comments:

104 phage from 8714 (a sequence) towards 3' end of gene

Trityl Group: Off Synthesis Scale: O.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8868 Thank you for using the HHMI Biopolymer Facility.

Ta = 60

A240 = ,330

30 2/970

Report for Synthetic Oligonucleotide . And the control of the control of

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8876

Product Information

Length: 20

Oligonucleotide Sequence: CAA GTA AGG GTA GCG GGC AC

Comments:

104 phage from primer 8848 towards 5' end of gene

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Mumber of Day(s) in the System: 2

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8876 Thank you for using the HHMI Biopolymer Facility.

A2Ga = 38.287

35 M/ 965 ml

Synthesis Request Sheet

FACILITY USE ONLY:

Sequence #

625 B

Column

Acceler C 407370 444444444

Oate Synthesized __ The state when the trade have stated and the state of the

Today's Date:

Sequence Length: 43

Requestor: P.I. Name:

Gersten LOWE

Synthesis Scale: 0.2 umole frityl Group: __On _X Off
Thio-Chemistry: __Yes _X No

Billing Dept.: HHMI

Phone Number: 7-4778

Account Number: 921099

SEQUENCE, 5'-> 3', IN TRIPLETS (* PLEASE START WITH A TRIPLET *)

GCG CGG ATC CTC ATC AAG CCT GGA ACC AGC ITT CAA GGT CTT C

User Comments:

104 phage PCR (for ab) from stop codon

Comments to Core Facility:

COLUMN 1 SET-UP VERSION 2.01

USER_NAME:

CYCLES USED:

SSCEAF3I- 1

ENDING METHOD:

Trityl OFF, Auto

ENDING PROCEDURE: DEPCRE03

SEQUENCE NAME: SEQUENCE LENGTH: 625B

DATE:

TIME:

COMMENT:

 $\sqrt{\text{5}$ - 606 066 ATC CTC ATC AAG CCT 66A ACC AGC TTT CAA

GGT CTT C -3

A260 = .662 19 al / 81 al

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8905

Product Information

Length: 20

Oligonucleotide Sequence: CAG GAA TTC AGG ATA TAA GG

Comments:

104 phage from primer 8868 towards 3' end of gene

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received: Date Synthesised:

Number of Day(s) in the System: 3

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8905 Thank you for using the HHMI Biopolymer Facility.

A260 = . 312

322/968

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8877

Product Information

Length: 20

Oligonucleotide Sequence: GGT AGT GCC ATG GTG ACC AA

Comments:

104 phage from primer 8848 towards 5' end of gene

Trityl Group: Off Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System: 2

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8877 Thank you for using the HHMI Biopolymer Facility.

A260 = .318

31 wl / 969 al

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8954

Product Information

Length: 20

Oligonucleotide Sequence: AGG TTG CAG ATG CAC CCT CT

Comments:

104 phage from primer 8905 towards 3' ut

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received:
Date Synthesised:

Number of Day(s) in the System: 4

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8954
Thank you for using the HHMI Biopolymer Facility.

A₂₆₀ = .272

37 l /963 ul

Date:

Investigator: LOWE

Individual User: Gersten

System Id:

8967

Product Information

the manufacture of the control of th

Length: 20

Oligonucleotide Sequence: GTA CTA GAG GGT GCA TCT GC

44 +

Comments:

104 phage from rp2 primer (sac insert) towards 5' end of gene

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

Date Received:

Date Synthesised:

Number of Day(s) in the System:

Ε,

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8967
Thank you for using the HHMI Biopolymer Facility.

Ta: SL

USE 104700 PTET9

 $A_{260} = .230$

442 /956

Report for Synthetic Oligonucleotide with dark was the same that th

Date:

System Id:

8713

Investigator: LOWE

Individual User: Gersten

Product Information

Length: 20

Oligonucleotide Sequence: ACC ACT CAA GCA AGG CTC TC

Comments:

104pstt7 towards rp2 primer

Trityl Group: Off

Synthesis Scale: 0.2 Micromole Scale

3

Date Received: Date Synthesised:

Number of Day(s) in the System:

If you have any problems, contact the Biopolymers Facility Your assigned System Id number is 8713 Thank you for using the HHMI Biopolymer Facility.

Azco = ,327

31 al digo 969 al 420